

CRF Errors Corrected by the STIC Systems Branch

Serial Number: 10/062,875A

CRF Processing Date: 7/9/02 0570
 Edited by: DC
 Verified by: DC (STIC staff) 0709
 #9

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/062,875A

DATE: 07/09/2002

TIME: 13:42:05

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\07092002\J062875A.raw

```

4 <110> APPLICANT: Cockerill, Franklin
5      Patel, Robin
6      Sloan, Lynne
8 <120> TITLE OF INVENTION: Detection of Bordetella
11 <130> FILE REFERENCE: 07039-253001
13 <140> CURRENT APPLICATION NUMBER: US 10/062,875A
14 <141> CURRENT FILING DATE: 2002-01-31
16 <150> PRIOR APPLICATION NUMBER: US 60/265,534
17 <151> PRIOR FILING DATE: 2001-01-31
19 <160> NUMBER OF SEQ ID NOS: 8
21 <170> SOFTWARE: FastSEQ for Windows Version 4.0
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 18
25 <212> TYPE: DNA
26 <213> ORGANISM: Artificial Sequence
W--> 27 <220> FEATURE:
28 <223> OTHER INFORMATION: Oligonucleotides
W--> 29 <400> SEQUENCE: 1
30 ccagttcctc aaggacgc 18
32 <210> SEQ ID NO: 2
33 <211> LENGTH: 23
34 <212> TYPE: DNA
35 <213> ORGANISM: Artificial Sequence
W--> 36 <220> FEATURE:
37 <223> OTHER INFORMATION: Oligonucleotides
W--> 38 <400> SEQUENCE: 2
39 gagttctggt aggtgtgagc gta 23
41 <210> SEQ ID NO: 3
42 <211> LENGTH: 28
43 <212> TYPE: DNA
44 <213> ORGANISM: Artificial Sequence
W--> 45 <220> FEATURE:
46 <223> OTHER INFORMATION: Oligonucleotides
W--> 47 <400> SEQUENCE: 3
48 caccgcttta cccgacctta ccgcccac 28
50 <210> SEQ ID NO: 4
51 <211> LENGTH: 28
52 <212> TYPE: DNA
53 <213> ORGANISM: Artificial Sequence
W--> 54 <220> FEATURE:
55 <223> OTHER INFORMATION: Oligonucleotides
W--> 56 <400> SEQUENCE: 4
57 gaccaatggc aaggccgaac gcttcac 28

```

RAW SEQUENCE LISTING

DATE: 07/09/2002

PATENT APPLICATION: US/10/062,875A

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Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\07092002\J062875A.raw

```

59 <210> SEQ ID NO: 5
60 <211> LENGTH: 18
61 <212> TYPE: DNA
62 <213> ORGANISM: Artificial Sequence
W--> 63 <220> FEATURE:
64 <223> OTHER INFORMATION: Oligonucleotides
W--> 65 <400> SEQUENCE: 5
66 ggcgatatca acgggtga 18
68 <210> SEQ ID NO: 6
69 <211> LENGTH: 19
70 <212> TYPE: DNA
71 <213> ORGANISM: Artificial Sequence
W--> 72 <220> FEATURE:
73 <223> OTHER INFORMATION: Oligonucleotides
W--> 74 <400> SEQUENCE: 6
75 cagggcaaac tcgtccatc 19
77 <210> SEQ ID NO: 7
78 <211> LENGTH: 24
79 <212> TYPE: DNA
80 <213> ORGANISM: Artificial Sequence
W--> 81 <220> FEATURE:
82 <223> OTHER INFORMATION: Oligonucleotides
W--> 83 <400> SEQUENCE: 7
84 gttcttcgaa ctgggttggc atac 24
86 <210> SEQ ID NO: 8
87 <211> LENGTH: 22
88 <212> TYPE: DNA
89 <213> ORGANISM: Artificial Sequence
W--> 90 <220> FEATURE:
91 <223> OTHER INFORMATION: Oligonucleotides
W--> 92 <400> SEQUENCE: 8
93 gtcaagacgc tggacaaggc tc 22

```

VERIFICATION SUMMARY

DATE: 07/09/2002

PATENT APPLICATION: US/10/062,875A

TIME: 13:42:06

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\07092002\J062875A.raw

L:27 M:283 W: Missing Blank Line separator, <220> field identifier
L:29 M:283 W: Missing Blank Line separator, <400> field identifier
L:36 M:283 W: Missing Blank Line separator, <220> field identifier
L:38 M:283 W: Missing Blank Line separator, <400> field identifier
L:45 M:283 W: Missing Blank Line separator, <220> field identifier
L:47 M:283 W: Missing Blank Line separator, <400> field identifier
L:54 M:283 W: Missing Blank Line separator, <220> field identifier
L:56 M:283 W: Missing Blank Line separator, <400> field identifier
L:63 M:283 W: Missing Blank Line separator, <220> field identifier
L:65 M:283 W: Missing Blank Line separator, <400> field identifier
L:72 M:283 W: Missing Blank Line separator, <220> field identifier
L:74 M:283 W: Missing Blank Line separator, <400> field identifier
L:81 M:283 W: Missing Blank Line separator, <220> field identifier
L:83 M:283 W: Missing Blank Line separator, <400> field identifier
L:90 M:283 W: Missing Blank Line separator, <220> field identifier
L:92 M:283 W: Missing Blank Line separator, <400> field identifier